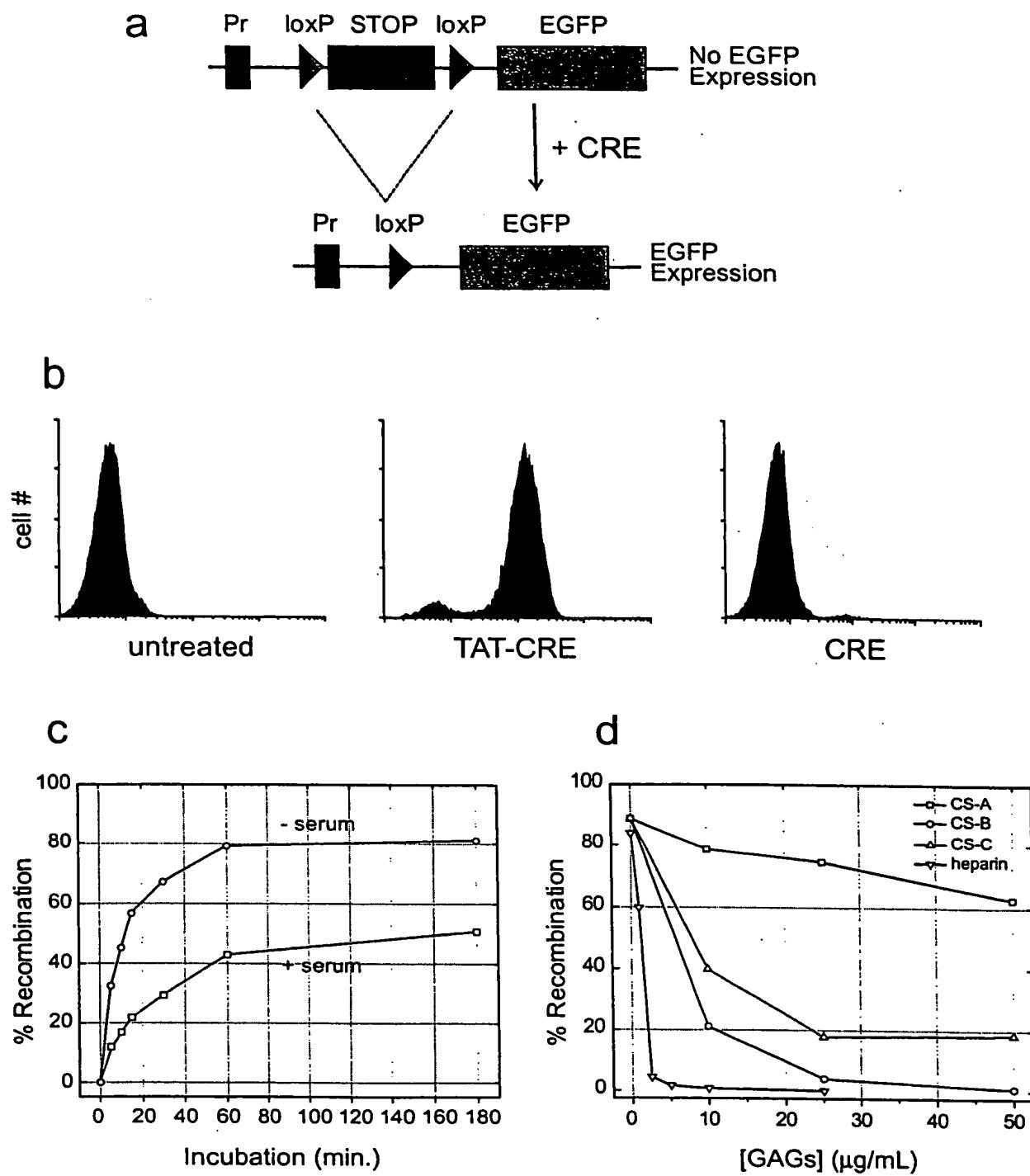
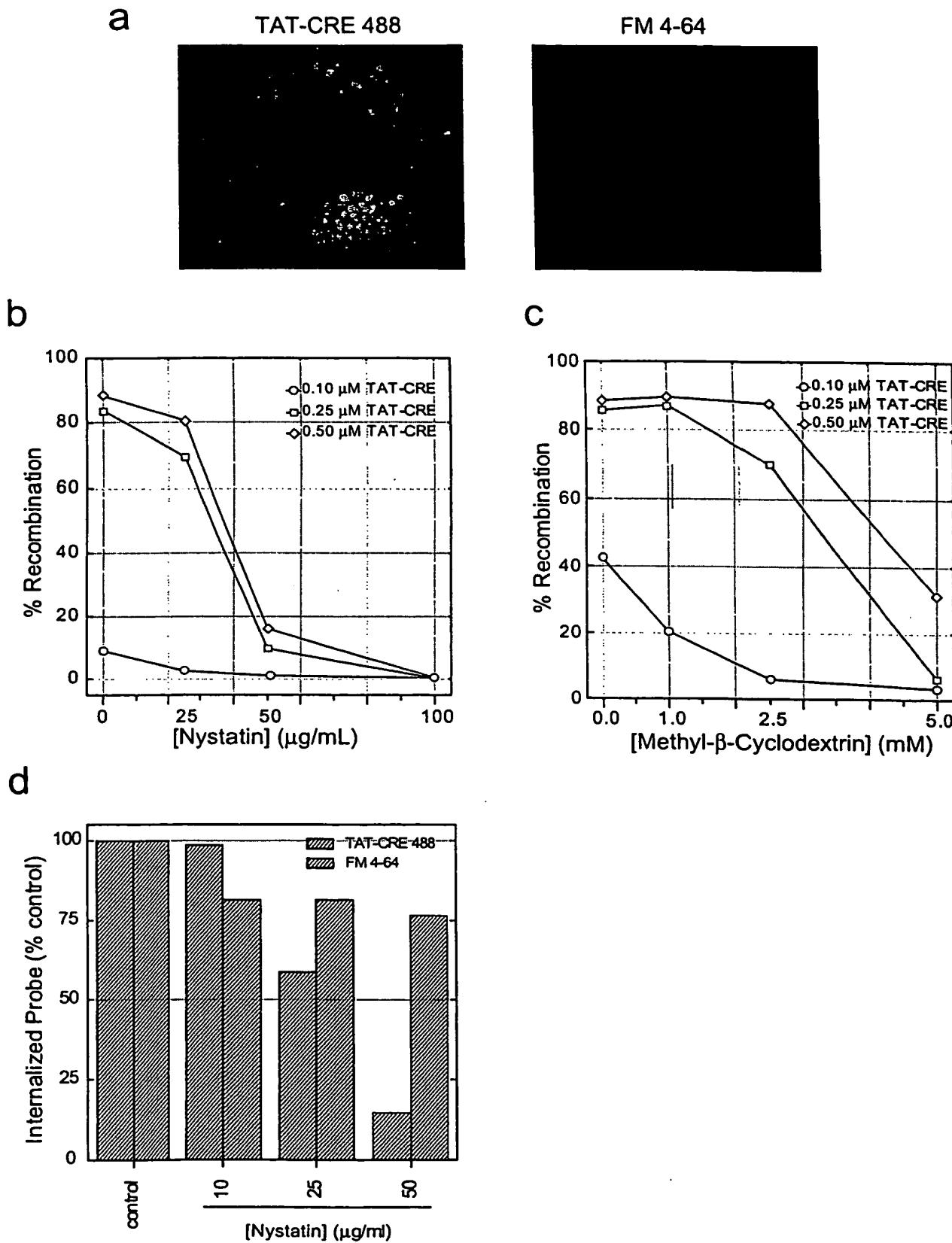
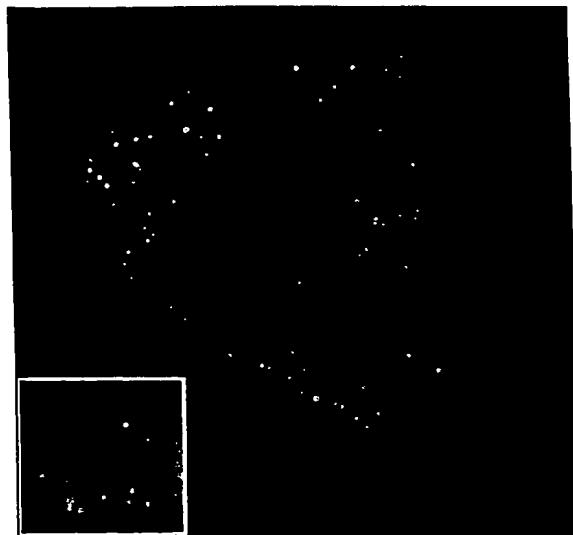


FIG. 1

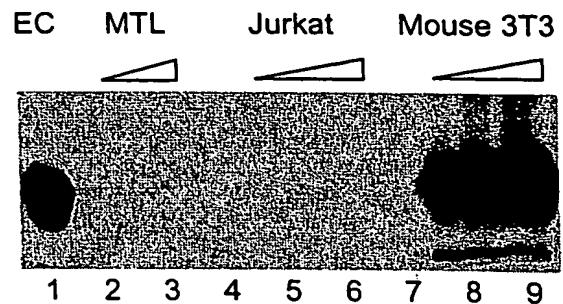
**FIG. 2A-D**

**FIG. 3A-D**

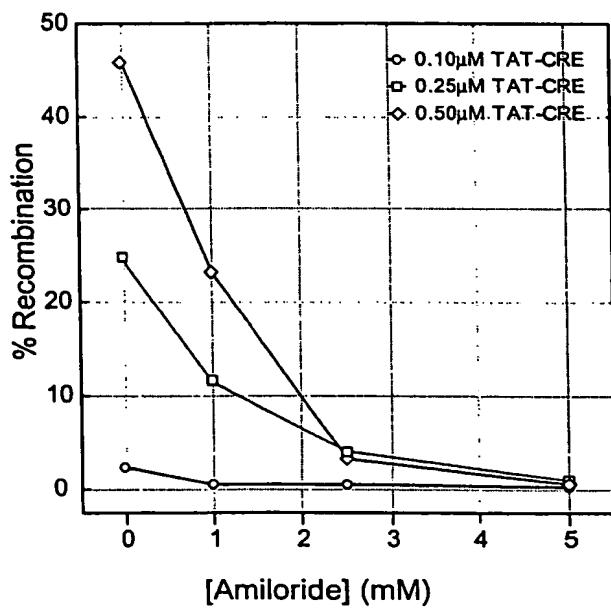
a



b



c



d

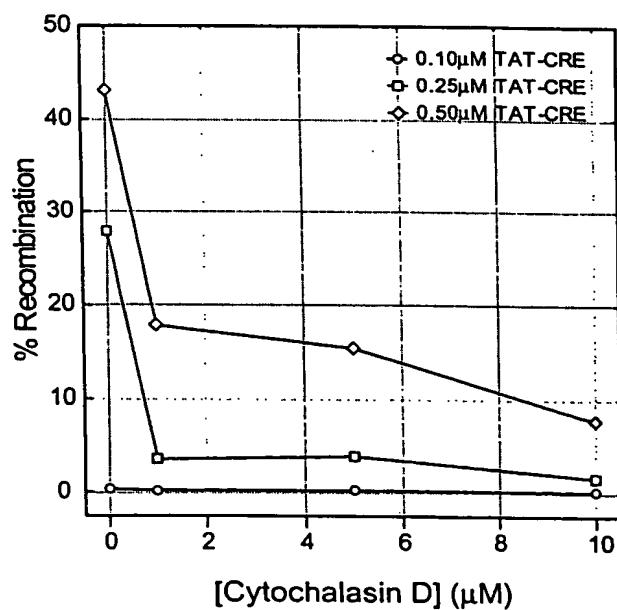


FIG. 4A-D

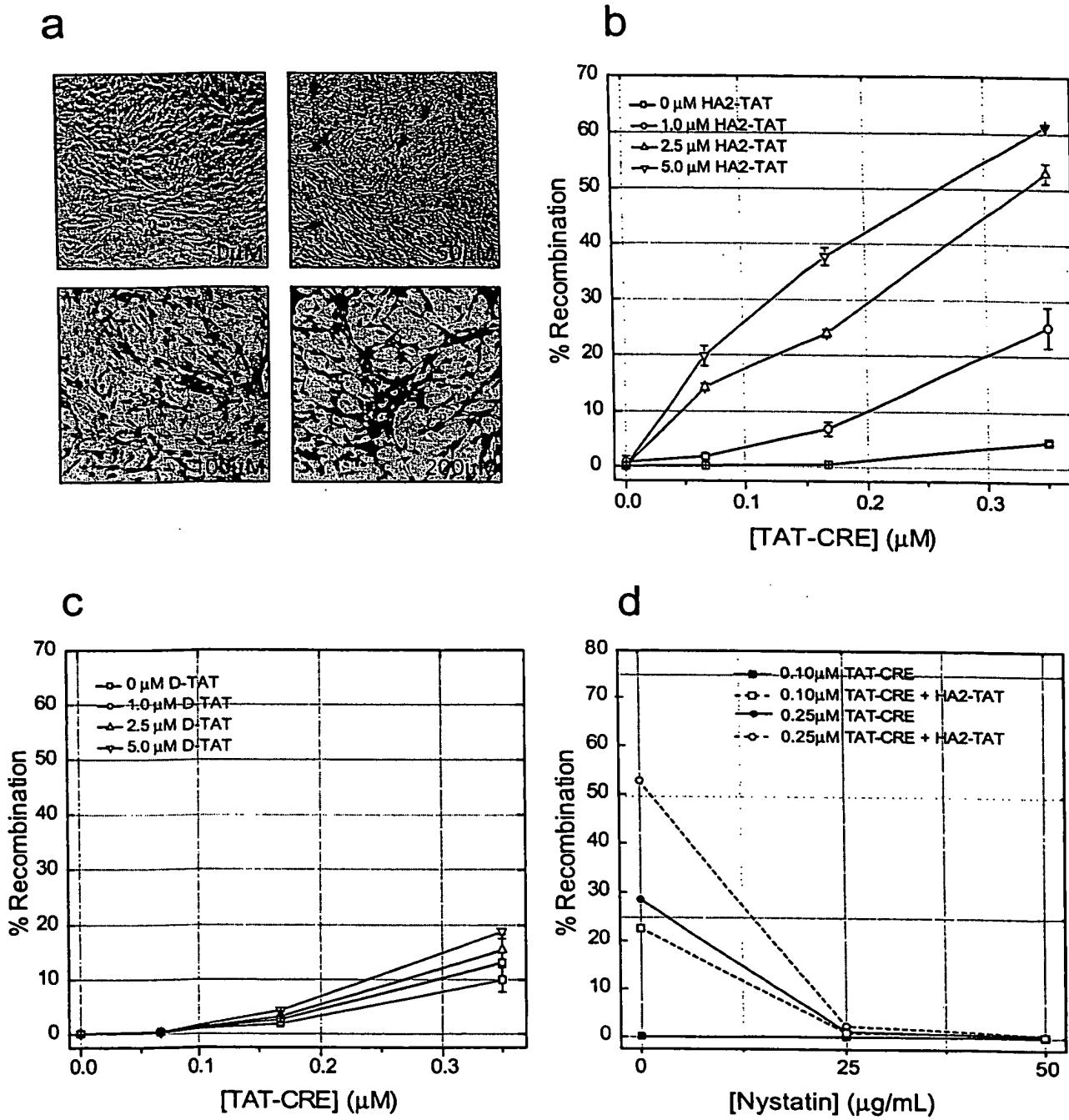
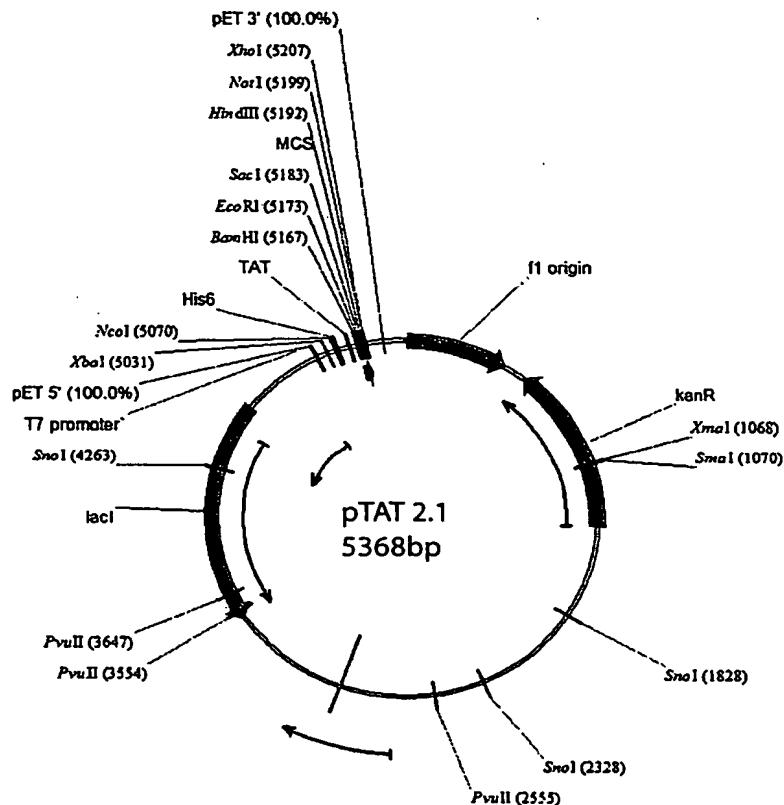


FIG. 5A-D



		T7 promoter													
		pET 5 (100.0x)													
4951	GGCTAGAGGA TCGAGATCTC GATCCGGCGA AATTATACCC ACTACATAA GGGGAAATGT GAGCGGATAA CAAITCCCCCT CTAGAAATAA CGCACTCTT AGCTCTAGAG CTAGGGCGCT TTAATTAAGC TGAATGATAA CCCCTTAACA CTCCGCTATT GTTAAGGGGA GATCTTAT														
	<u>Nco</u> <u>M</u> G S S H H H H H H S S G L V P R G S H														
5041	TTTGTAACTT CTTARGAAAG GAGATACCC AIGGGCAGCA GGCATCACTCA TCACTCATCAC AGCACGGCC TGGTCGCGCG CGGCAGCCAT AAAACAAATT GAAATCTTC CTCTATATGG TACCCGTCTG CCGTAGATGT AGTAGTGTG TCGTCGCGCG ACCACGGCGC GCGCTCGGTAA	<u>TAT</u> <u>MCS</u>													
	<u>EcoRI</u> <u>BamH</u> M R K K R R Q R R R G S D P N E S S V D K L A A A L E H H H H														
5131	ATGAGGAAGA AGCCGGAGACG GCGACGAAGA GGCTCGGATC CGAATTCGGAG CTCCGTCGAC AAAGCTTCGG CGCGACTCTGA GCACCAACAC TACTCTCTCTG TCGGCTCTGT CGCTGCTCTCT CGGAGGCTAG CTCTAACCTC GAGGCCAGCTG TTCCGAAACGCC GCGGTGAGCT CGTGGTGGTGT	<u>Sac</u> <u>Nde</u> <u>Kpn</u>													
	<u>H H H</u>														
5221	CACCAACCACT GAGATCCGGC TCTTAACAAA GCGCGAARRGG AAAGCTGAGTT GGCTGCTGCC ACCGCTGAGC AATAACTAGC ATAAACCCCTI GTGGTGGTGA CTCTAGGCCG AGCAATGTTT CGGGCTTICC TTCCGACTCAA CGGACGACGG TGGCGACTCG TTATTGATCG TATTGGGGAA	<u>pET 5 (100.0x)</u>													
	<u>GGGGCTCTTAA</u> AACGGGTCTT GAGGGTTTT TCTCTGAAAG GAGGAACATAA AICCGGAT CCCCGGAGAT TTGCCCCAGAA CTCCCCAAA AACGACTTTC CTCCCTGATA TAGGGCTA														

FIG. 6

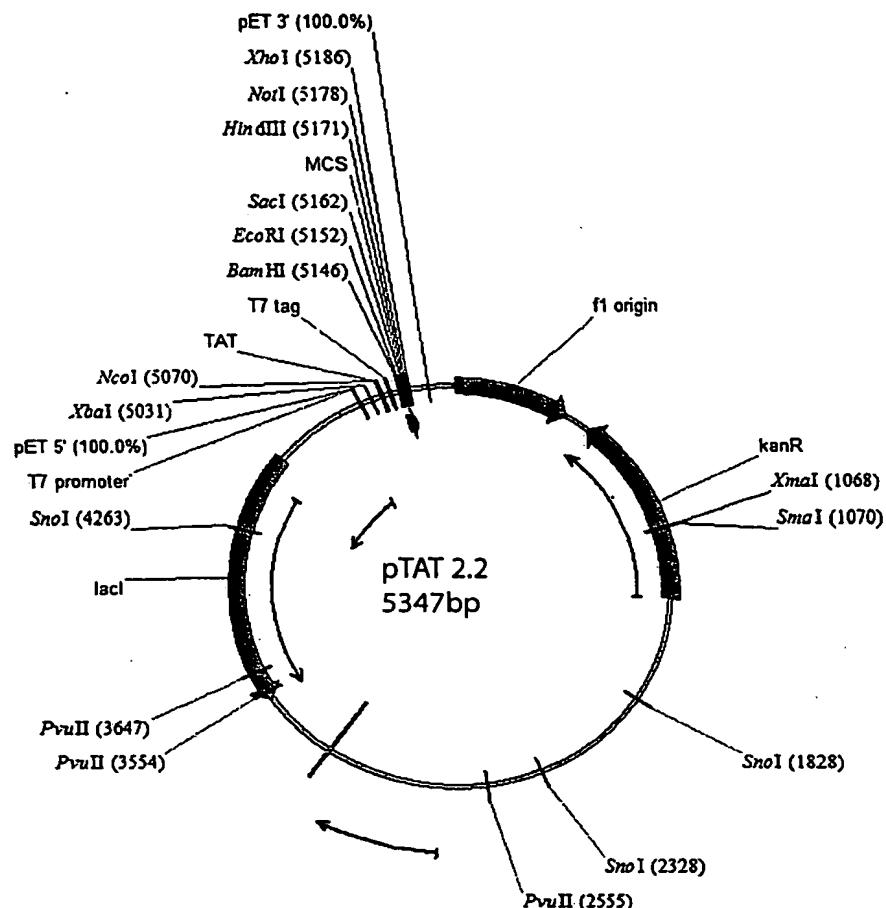


FIG. 7

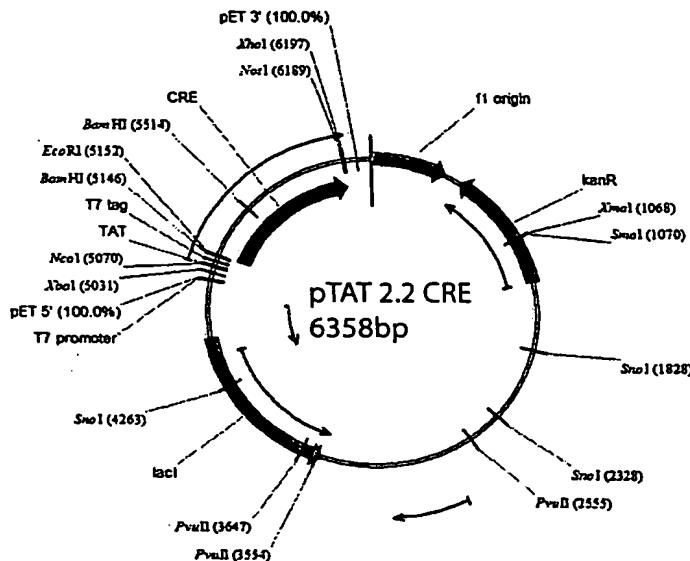


FIG. 8